

1600

OIEP

1642 #5

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/944,449

DATE: 12/10/2001  
TIME: 15:44:09

Input Set : N:\paola\09944449.txt  
Output Set: N:\CRF3\12102001\I944449.raw

RECEIVED

JAN 18 2002

TECH CENTER 1600/2900

ENTERED

3 <110> APPLICANT: Baker, Kevin  
4 Botstein, David  
5 Eaton, Dan  
6 Ferrara, Napoleone  
7 Filvaroff, Ellen  
8 Gerritsen, Mary  
9 Goddard, Audrey  
10 Godowski, Paul  
11 Grimaldi, Christopher  
12 Gurney, Austin  
13 Hillan, Kenneth  
14 Kljavin, Ivar  
15 Napier, Mary  
16 Roy, Margaret  
17 Tumas, Daniel  
18 Wood, William  
20 <120> TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC  
21 ACIDS ENCODING THE SAME  
23 <130> FILE REFERENCE: P2548P1C1  
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25 <141> CURRENT FILING DATE: 2001-08-30  
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28 <151> PRIOR FILING DATE: 2001-05-25  
30 <150> PRIOR APPLICATION NUMBER: 60/069,334  
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33 <150> PRIOR APPLICATION NUMBER: 60/069335  
W--> 34 <151> PRIOR FILING DATE: December 11, 1997  
36 <150> PRIOR APPLICATION NUMBER: 60/069,278  
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45 <150> PRIOR APPLICATION NUMBER: 60/069,694  
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48 <150> PRIOR APPLICATION NUMBER: 60/069,702  
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54 <150> PRIOR APPLICATION NUMBER: 60/069,873  
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57 <150> PRIOR APPLICATION NUMBER: 60/068,017  
W--> 58 <151> PRIOR FILING DATE: December 18, 1997  
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W--> 61 <151> PRIOR FILING DATE: January 5, 1998  
63 <150> PRIOR APPLICATION NUMBER: 60/074,086  
W--> 64 <151> PRIOR FILING DATE: February 9, 1998

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78 <150> PRIOR APPLICATION NUMBER: 60/146,222  
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84 <150> PRIOR APPLICATION NUMBER: PCT/US98/25108  
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90 <150> PRIOR APPLICATION NUMBER: 09/218,517  
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93 <150> PRIOR APPLICATION NUMBER: 09/254,311  
W--> 94 <151> PRIOR FILING DATE: March 3, 1999  
96 <150> PRIOR APPLICATION NUMBER: PCT/US99/12252  
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99 <150> PRIOR APPLICATION NUMBER: PCT/US99/21090  
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102 <150> PRIOR APPLICATION NUMBER: PCT/US99/28409  
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132 <150> PRIOR APPLICATION NUMBER: PCT/US00/32678  
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141 <211> LENGTH: 2454
142 <212> TYPE: DNA
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148 caccaggact gtgttgaagg gtgttttttt tctttttaa gtaatacctc 100
150 ctcactctttt cttctttacac agtgtctgag aacattttaca ttatagataa 150
152 gtagtacatg gtggataact tctactttta ggaggactac tctcttctga 200
154 cagtcctaga ctggtcttct acactaagac accatgaagg agtatgtgct 250
156 cctattattc ctggctttgt gctctgccaa acccttcttt agcccttcac 300
158 acatcgcact gaagaatatg atgctgaagg atatggaaga cacagatgat 350
160 gatgatgatg atgatgatga tgatgatgat gatgaggaca actctctttt 400
162 tccaacaaga gagccaagaa gccatttttt tccatttgat ctgtttccaa 450
164 tgtgtccatt tggatgtcag tgctattcac gagttgtaca ttgctcagat 500
166 ttaggtttga cctcagtcac aaccaacatt ccatttgata ctcgaatgct 550
168 tgatcttcaa aacaataaaa ttaaggaaat caaagaaaat gattttaaag 600
170 gactcacttc actttatggt ctgatcctga acaacaacaa gctaacgaag 650
172 attcacccaa aagcctttct aaccacaaag aagttgcgaa ggctgtatct 700
174 gtcccacaat caactaagtg aaataccact taatcttccc aaatcattag 750
176 cagaactcag aattcatgaa aataaagtta agaaaataca aaaggacaca 800
178 ttcaaaggaa tgaatgcttt acacgttttg gaaatgagtg caaacctctc 850
180 tgataataat gggatagagc caggggcatt tgaagggtg acggtgttcc 900
182 atatcagaat tgcagaagca aaactgacct cagttcctaa aggcttacca 950
184 ccaactttat tggagcttca cttagattat aataaaattt caacagtgga 1000
186 acttgaggat tttaaacgat acaaagaact acaaaggctg ggcctaggaa 1050
188 acaacaaaat cacagatata gaaaatggga gtcttgctaa cataccacgt 1100
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192 attaccagag ttgaaatacc tccagataat ctcccttcat tctaattcaa 1200
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196 aaatctttat acagtgcaat aagtttattc aacaaccggt tgaaatactg 1300
198 ggaaatgcaa cctgcaacat ttcgttgtgt tttgagcaga atgagtgttc 1350
200 agcttgaggaa ctttggaatg taataattag taattggtaa tgtccattta 1400
202 atataagatt caaaaatccc tacatttgga atacttgaac tctattaata 1450
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206 actgacttat tttatgacaa gaaatttcaa cggaattttg ccaaactatt 1550
208 gatacataag gggttgagag aaacaagcat ctattgcagt ttcctttttg 1600
210 cgtacaaaat atcttacata aatctcatgc ttgaccattc ctttcttcat 1650
212 aacaaaaaag taagatatc ggtatttaac actttgttat caagcacatt 1700
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224 taaagccttc agtaaatgtt cattaccaac ttgataaatg ctactcataa 2000
226 gagctggttt ggggctatag catatgcttt ttttttttta attattacct 2050
228 gattttaaaa tctctgtaaa aacgtgtagt gtttcataaa atctgtaact 2100
230 cgcattttta tgatccgcta ttataagctt ttaatagcat gaaaattgtt 2150
232 aggctatata acattgccac ttcaactcta aggaatattt ttgagatata 2200

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234 cctttggaag accttgcttg gaagagcctg gacactaaca attctacacc 2250
236 aaattgtctc ttcaaatacg tatggactgg ataactctga gaaacacatc 2300
238 tagtataact gaataagcag agcatcaaat taaacagaca gaaaccgaaa 2350
240 gctctatata aatgctcaga gttctttatg tattttcttat tggcattcaa 2400
242 catatgtaaa atcagaaaaac agggaaattt tcattaaaaa tattggtttg 2450
244 aaat 2454
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247 <211> LENGTH: 379
248 <212> TYPE: PRT
249 <213> ORGANISM: Homo Sapien
251 <400> SEQUENCE: 2
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253 1 5 10 15
255 Lys Pro Phe Phe Ser Pro Ser His Ile Ala Leu Lys Asn Met Met
256 20 25 30
258 Leu Lys Asp Met Glu Asp Thr Asp Asp Asp Asp Asp Asp Asp
259 35 40 45
261 Asp Asp Asp Asp Asp Glu Asp Asn Ser Leu Phe Pro Thr Arg Glu
262 50 55 60
264 Pro Arg Ser His Phe Phe Pro Phe Asp Leu Phe Pro Met Cys Pro
265 65 70 75
267 Phe Gly Cys Gln Cys Tyr Ser Arg Val Val His Cys Ser Asp Leu
268 80 85 90
270 Gly Leu Thr Ser Val Pro Thr Asn Ile Pro Phe Asp Thr Arg Met
271 95 100 105
273 Leu Asp Leu Gln Asn Asn Lys Ile Lys Glu Ile Lys Glu Asn Asp
274 110 115 120
276 Phe Lys Gly Leu Thr Ser Leu Tyr Gly Leu Ile Leu Asn Asn Asn
277 125 130 135
279 Lys Leu Thr Lys Ile His Pro Lys Ala Phe Leu Thr Thr Lys Lys
280 140 145 150
282 Leu Arg Arg Leu Tyr Leu Ser His Asn Gln Leu Ser Glu Ile Pro
283 155 160 165
285 Leu Asn Leu Pro Lys Ser Leu Ala Glu Leu Arg Ile His Glu Asn
286 170 175 180
288 Lys Val Lys Lys Ile Gln Lys Asp Thr Phe Lys Gly Met Asn Ala
289 185 190 195
291 Leu His Val Leu Glu Met Ser Ala Asn Pro Leu Asp Asn Asn Gly
292 200 205 210
294 Ile Glu Pro Gly Ala Phe Glu Gly Val Thr Val Phe His Ile Arg
295 215 220 225
297 Ile Ala Glu Ala Lys Leu Thr Ser Val Pro Lys Gly Leu Pro Pro
298 230 235 240
300 Thr Leu Leu Glu Leu His Leu Asp Tyr Asn Lys Ile Ser Thr Val
301 245 250 255
303 Glu Leu Glu Asp Phe Lys Arg Tyr Lys Glu Leu Gln Arg Leu Gly
304 260 265 270
306 Leu Gly Asn Asn Lys Ile Thr Asp Ile Glu Asn Gly Ser Leu Ala
307 275 280 285

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309 Asn Ile Pro Arg Val Arg Glu Ile His Leu Glu Asn Asn Lys Leu
310                290                295                300
312 Lys Lys Ile Pro Ser Gly Leu Pro Glu Leu Lys Tyr Leu Gln Ile
313                305                310                315
315 Ile Phe Leu His Ser Asn Ser Ile Ala Arg Val Gly Val Asn Asp
316                320                325                330
318 Phe Cys Pro Thr Val Pro Lys Met Lys Lys Ser Leu Tyr Ser Ala
319                335                340                345
321 Ile Ser Leu Phe Asn Asn Pro Val Lys Tyr Trp Glu Met Gln Pro
322                350                355                360
324 Ala Thr Phe Arg Cys Val Leu Ser Arg Met Ser Val Gln Leu Gly
325                365                370                375
327 Asn Phe Gly Met
330 <210> SEQ ID NO: 3
331 <211> LENGTH: 20
332 <212> TYPE: DNA
333 <213> ORGANISM: Artificial Sequence
335 <220> FEATURE:
336 <223> OTHER INFORMATION: Synthetic Oligonucleotide Probe
338 <400> SEQUENCE: 3
339 ggaaatgagt gcaaaccctc 20
341 <210> SEQ ID NO: 4
342 <211> LENGTH: 24
343 <212> TYPE: DNA
344 <213> ORGANISM: Artificial Sequence
346 <220> FEATURE:
347 <223> OTHER INFORMATION: Synthetic Oligonucleotide Probe
349 <400> SEQUENCE: 4
350 tcccaagctg aacactcatt ctgc 24
352 <210> SEQ ID NO: 5
353 <211> LENGTH: 50
354 <212> TYPE: DNA
355 <213> ORGANISM: Artificial Sequence
357 <220> FEATURE:
358 <223> OTHER INFORMATION: Synthetic Oligonucleotide Probe
360 <400> SEQUENCE: 5
361 gggtgacggt gttccatc agaatgcag aagcaaaact gacctcagtt 50
363 <210> SEQ ID NO: 6
364 <211> LENGTH: 3441
365 <212> TYPE: DNA
366 <213> ORGANISM: Homo Sapien
368 <400> SEQUENCE: 6
369 cggacgcgtg ggcggacgcg tgggcccgcg gcaccgcccc cggcccggcc 50
371 ctccgccctc cgcactcgcg cctccctccc tccgcccgcg cccgcgccct 100
373 cctccctccc tcctccccag ctgtcccgtt cgcgtcatgc cgagcctccc 150
375 ggccccgcgc gcccgctgc tgcctctcgg gctgctgctg ctcggtccc 200
377 ggccgcgcgc cggcgccggc ccagagcccc ccgtgctgcc catccgttct 250
379 gagaaggagc cgtgcccgt tcggggagcg gcaggctgca ccttcggcgg 300
381 gaaggtctat gccttggacg agacgtggca cccggaccta gggcagccat 350

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## VERIFICATION SUMMARY

PATENT APPLICATION: US/09/944,449

DATE: 12/10/2001

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Output Set: N:\CRF3\12102001\I944449.raw

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